## **AMENDMENTS TO THE SPECIFICATION**

On page 13 of the specification, please replace the paragraph spanning lines 4 to 8 with the following amended paragraph:

The SLR method does not depend on the presence of a high level of homology between polynucleotides to be shuffled. The invention can be used to non-stochastically generate libraries (or sets) of progeny molecules comprised of over  $\underline{10^{100}}$  [[10100]] different chimeras. Conceivably, SLR can even be used to generate libraries comprised of over  $\underline{10^{1000}}$  [[101000]] different progeny chimeras.